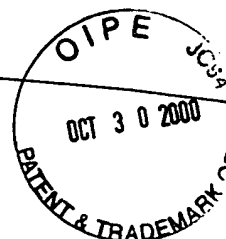


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## SEQUENCE LISTING



TECH CENTER 1600/2900

<110> Keyt, Bruce A..  
 Nguyen, Francis H.  
 Ferrara, Napoleone

<120> NUCLEIC ACIDS ENCODING VARIANTS OF VASCULAR ENDOTHELIAL  
 CELL GROWTH FACTOR

<130> A-62326-2

<140> 09/346,069

<141> 1999-07-01

<150> 08/567,200

<151> 1995-12-05

<150> 60/002,827

<151> 1995-08-25

<160> 42

<170> PatentIn Ver. 2.1

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Met

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Asn Phe Leu Leu Ser Trp Val His Trp Ser Leu Ala Leu Leu Leu Tyr

5

10

15

ctc cac cat gcc aag tgg tcc cag gct gca ccc atg gca gaa gga gga 155

Leu His His Ala Lys Trp Ser Gln Ala Ala Pro Met Ala Glu Gly Gly

20

25

30

ggg cag aat cat cac gaa gtg gtg aag ttc atg gat gtc tat cag cgc 203

Gly Gln Asn His His Glu Val Val Lys Phe Met Asp Val Tyr Gln Arg

35

40

45

agc tac tgc cat cca atc gag acc ctg gtg gac atc ttc cag gag tac 251  
 Ser Tyr Cys His Pro Ile Glu Thr Leu Val Asp Ile Phe Gln Glu Tyr  
 50 55 60 65

cct gat gag atc gag tac atc ttc aag cca tcc tgt gtg ccc ctg atg 299  
 Pro Asp Glu Ile Glu Tyr Ile Phe Lys Pro Ser Cys Val Pro Leu Met  
 70 75 80

cga tgc ggg ggc tgc tgc aat gac gag ggc ctg gag tgt gtg ccc act 347  
 Arg Cys Gly Gly Cys Cys Asn Asp Glu Gly Leu Glu Cys Val Pro Thr  
 85 90 95

gag gag tcc aac atc acc atg cag att atg cgg atc aaa cct cac caa 395  
 Glu Glu Ser Asn Ile Thr Met Gln Ile Met Arg Ile Lys Pro His Gln  
 100 105 110

ggc cag cac ata gga gag atg agc ttc cta cag cac aac aaa tgt gaa 443  
 Gly Gln His Ile Gly Glu Met Ser Phe Leu Gln His Asn Lys Cys Glu  
 115 120 125

tgc aga cca aag aaa gat aga gca aga caa gaa aat ccc tgt ggg cct 491  
 Cys Arg Pro Lys Lys Asp Arg Ala Arg Gln Glu Asn Pro Cys Gly Pro  
 130 135 140 145

tgc tca gag cgg aga aag cat ttg ttt gta caa gat ccg cag acg tgt 539  
 Cys Ser Glu Arg Arg Lys His Leu Phe Val Gln Asp Pro Gln Thr Cys  
 150 155 160

aaa tgt tcc tgc aaa aac aca gac tgc cgt tgc aag gcg agg cag ctt 587  
 Lys Cys Ser Cys Lys Asn Thr Asp Ser Arg Cys Lys Ala Arg Gln Leu  
 165 170 175

gag tta aac gaa cgt act tgc aga tgt gac aag ccg agg cgg tga 632  
 Glu Leu Asn Glu Arg Thr Cys Arg Cys Asp Lys Pro Arg Arg  
 180 185 190

gccggggcagg aggaaggagc ctccctcagg gtttcgggaa ccagatctct caccaggaaa 692

gactgataca gaacgatcga tacagaaacc acgctgccgc caccacacca tcaccatcga 752

cagaacagtc cttaatccag aaacctgaaa tgaaggaaga ggagactctg cgcagagcac 812

tttgggtccg gagggcgaga ctccggcgga agcattcccg ggcgggtgac ccagcacggt 872

ccctcttggga attggattcg ccattttatt tttcttgctg ctaaatacacc gagcccgga 932

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<210> 2  
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1 5 10 15  
Tyr Leu His His Ala Lys Trp Ser Gln Ala Ala Pro Met Ala Glu Gly  
20 25 30  
Gly Gly Gln Asn His His Glu Val Val Lys Phe Met Asp Val Tyr Gln  
35 40 45  
Arg Ser Tyr Cys His Pro Ile Glu Thr Leu Val Asp Ile Phe Gln Glu  
50 55 60  
Tyr Pro Asp Glu Ile Glu Tyr Ile Phe Lys Pro Ser Cys Val Pro Leu  
65 70 75 80  
Met Arg Cys Gly Gly Cys Cys Asn Asp Glu Gly Leu Glu Cys Val Pro  
85 90 95  
Thr Glu Glu Ser Asn Ile Thr Met Gln Ile Met Arg Ile Lys Pro His  
100 105 110  
Gln Gly Gln His Ile Gly Glu Met Ser Phe Leu Gln His Asn Lys Cys  
115 120 125  
Glu Cys Arg Pro Lys Lys Asp Arg Ala Arg Gln Glu Asn Pro Cys Gly  
130 135 140  
Pro Cys Ser Glu Arg Arg Lys His Leu Phe Val Gln Asp Pro Gln Thr  
145 150 155 160  
Cys Lys Cys Ser Cys Lys Asn Thr Asp Ser Arg Cys Lys Ala Arg Gln  
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<210> 8  
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<400> 8  
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<210> 11  
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<210> 12  
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<210> 14  
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<210> 15  
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<400> 15  
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<210> 16  
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<212> DNA  
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<210> 17  
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<220>  
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<210> 19  
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<400> 19  
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<400> 20  
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<210> 24  
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<400> 24  
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<210> 25  
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<212> DNA  
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<210> 30  
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<400> 30  
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<210> 31  
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<210> 32

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<210> 33

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<210> 34

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<210> 35

<211> 30

<212> DNA

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<210> 40

<211> 27

<212> DNA

<213> Artificial Sequence

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<210> 41

<211> 27

<212> DNA

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<210> 42

<211> 27

<212> DNA

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<220>

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Concluded